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Sequence Listing could not be accepted.
If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).
Reviewer: Anne Corrigan
Timestamp: Mon May 07 11:05:48 EDT 2007
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The <213> response is invalid, per 1.823 of Sequence Rules.  The only
valid <213> responses are "Artificial Sequence," "Unknown," or the
Genus/species.  Same error in sequences 7-11.
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Application No: 10580901   Version No: 1.0

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       Berry, Carole
       Thomas, Mark
       Kambadur, Ravi
       Bower, Robert Syndecombe

<120>  Novel Muscle Growth Regulator

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aaacaagaat atagtcgtta tcagaggtgg agacatttag aagttgttct taatcagagt     300

gaagcttgta cttcggaaag tcagcctcac tcctcagcac tcacagcacc tagttctcca     360

ggttcctcct ggatgaaaaa ggaccagccc acctttaccc tccgacaagt tggaataata     420

tgtgagcgtc tcttaaaaga ctatgaagat aaaattcggg aggaatatga gcaaatcctc     480

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Gly Pro Thr Pro Gly Leu Arg Pro Pro Asp Ala Glu Pro Pro Leu
            35   40   45

Leu Gln Thr Gln Thr Pro Pro Pro Thr Leu Gln Gln Pro Ala Pro Pro
    50   55   60

Gly Ser Glu Arg Arg Leu Pro Thr Pro Glu Gln Ile Phe Gln Asn Ile
65   70   75   80

Lys Gln Glu Tyr Ser Arg Tyr Gln Arg Trp Arg His Leu Glu Val Val
                85   90   95

Leu Asn Gln Ser Glu Ala Cys Thr Ser Glu Ser Gln Pro His Ser Ser
            100   105   110

Ala Leu Thr Ala Pro Ser Ser Pro Gly Ser Ser Trp Met Lys Lys Asp
        115   120   125

Gln Pro Thr Phe Thr Leu Arg Gln Val Gly Ile Ile Cys Glu Arg Leu
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Leu Lys Asp Tyr Glu Asp Lys Ile Arg Glu Glu Tyr Glu Gln Ile Leu
145   150   155   160

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            35   40   45

Leu Gln Thr Gln Ile Pro Pro Thr Leu Gln Gln Pro Ala Pro Pro
    50   55   60

Gly Ser Asp Arg Arg Leu Pro Thr Pro Glu Gln Ile Phe Gln Asn Ile
65   70   75   80

Lys Gln Glu Tyr Ser Arg Tyr Gln Arg Trp Arg His Leu Glu Val Val
                85   90   95

Leu Asn Gln Ser Glu Ala Cys Thr Ser Glu Ser Gln Pro His Ser Ser
            100   105   110

Thr Leu Thr Ala Pro Ser Ser Pro Gly Ser Ser Trp Met Lys Lys Asp
            115   120   125

Gln Pro Thr Phe Thr Leu Arg Gln Val Gly Ile Ile Cys Glu Arg Leu
    130   135   140

Leu Lys Asp Tyr Glu Asp Lys Ile Arg Glu Glu Tyr Glu Gln Ile Leu
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